

FT	Peptide	1..28
FT		/note= "signal peptide"
FT	Protein	29..666
FT		/note= "mature protein"
XX		
PN	MO9827805-A1.	
XX		
PD	02-JUL-1998.	
XX		
PE	22-DEC-1997;	97WO-AU00874.
XX		
PR	20-DEC-1996;	96AU-0004275.
XX		
PA	(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.	
XX		
PI	Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;	
XX		
DR	WPI: 1998-377279/32.	
DR	N-PSDB: V42311.	
XX		
PT	Novel anti-microbial protein from e.g. Macadamia integrifolia -	
PT	useful for controlling microbial infestations of plants or mammals	
XX		
PS	Claim 1; Page 39-41; 96pp; English.	
XX		
CC	The sequence is that of an antimicrobial protein which can	
CC	be used to control microbial infestations in plants and mammalian	
CC	animals.	
XX		
SO	Sequence 666 AA;	

```

Query Match      27.5%; Score 105; DB 19; Length 666;
Best Local Similarity 34.3%; Pred. No. 6,4e-05;
Matches 23; Conservative 15; Mismatches 25; Indels 4; Gaps

QY          3 DEDDRRG--GSHSLQCVQRCHQERPRYSHARCVAQECDQDQOQHGR-HEQEEEGRGRGMH 59
             :|::||           |:|||::||   ::||::||::|||::||   |||::
Db    182 eednkrdpqgqreyedcrrrceqqeqprqgy-qqrccrqeqrgdqrgdllnpqrgsgry 240
              |||::
QY          60 GEGERE 66
Db    241 eegeekq 247

RESULT       5
ID    W62832 standard; Protein; 590 AA.
XX    AC    W62832;
XX    DT    27-OCT-1998 (first entry)
XX    DE    Gossypium hirsutum antimicrobial protein.
XX    KW    antimicrobial protein; infestation; control.
XX    OS    Gossypium hirsutum.
XX    PN    WO9827805-A1.
XX    PD    02-JUL-1998.
XX    PF    22-DEC-1997; 97WO-AU00874.
XX    PR    20-DEC-1996; 96AU-0004275.
XX    PA    (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
XX    PI    Bower NI, Goulter KC, Green JL, Mannens JM, Marcus JP;
XX    WP    1998-377279/32.
XX    DR

```

PT Novel anti-microbial protein from e.g. *Macadamia integrifolia* -
 PT useful for controlling microbial infestations of plants or mammals
 XX
 XX
 PS Claim 1; Page 49-51; 96pp; English.
 PS
 XX
 CC The sequence is that of an antimicrobial protein which can
 CC be used to control microbial infestations in plants and mammalian
 CC animals.
 XX
 XX Sequence 590 AA;
 SO

Query Match	24.3%	Score 93	DB 19	Length 590
Best Local Similarity	29.4%	Pred. 0.0016		
Matches	20	Conservative	16	Mismatches 24
				Indels 8
				Gaps 3
QY	3	DEDDRGRGSHLOOCVORCRORERPR-YSHARCVOEGRDQOOH---	GRHGEFEGRGRGCM	58
	::: ::	::	::	
Db	114	egeqgqsgqrqfdecqghcqhgcqqrpekkkqcyrecrekqyenpwtgereteeaeete--		171
QY	59	HGEGEREE	66	
		::		
Db	172	--egeqeq	177	
RESULT	6			
W62830				
ID	W62830	standard	Protein	625 AA.

Query Match	23.88;	Score 91;	DB 19;	Length 625;
DT	27-OCT-1998	(first entry)		
DE	Macadamia integrifolia	antimicrobial protein.		
XX				
XX	antimicrobial protein;	infestation; control;		
KW				
XX				
OS	Macadamia integrifolia.			
XX				
FH	Key	location/Qualifiers		
FT	Peptide	1..28		
FT		/note="signal peptide"		
FT	Protein	29..666		
FT		/note="mature protein"		
XX				
PN	W09827805-A1.			
XX				
PD	02-JUL-1998.			
XX				
PF	22-DEC-1997;	97WO-AU00874.		
XX				
PR	20-DEC-1996;	96AU-0004275.		
XX				
PA	(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.			
XX				
PI	Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;			
XX				
DR	WP1: 1998-377279/32.			
DR	N-PSDB: V42316.			
XX				
PT	Novel anti-microbial protein from e.g. Macadamia integrifolia -			
PT	useful for controlling microbial infestations of plants or mammals			
PS	Claim 1; Page 43-45; 96pp; English.			
XX				
CC	The sequence is that of an antimicrobial protein which can			
CC	be used to control microbial infestations in plants and mammalian			
CC	animals.			
XX				
XX	Sequence	625 AA;		

XX	Bacillus subtilis.	
OS		
XX	Key	Location/Qualifiers
FT	Misc-difference 67	/note= "Encoded by TAA stop codon"
FT	Misc-difference 91	/note= "Encoded by TAA stop codon"
FT	Misc-difference 92	/note= "Encoded by TAA stop codon"
FT	Misc-difference 111	/note= "Encoded by TAA stop codon"
FT	Misc-difference 120	/note= "Encoded by TGA stop codon"
FT	Misc-difference 133	/note= "Encoded by TAG stop codon"
FT	Misc-difference 146	/note= "Encoded by TAG stop codon"
FT	Misc-difference 151	/note= "Encoded by TAA stop codon"
FT	Misc-difference 176	/note= "Encoded by TGA stop codon"
FT	Misc-difference 184	/note= "Encoded by TGA stop codon"
FT	Misc-difference 201	/note= "Encoded by TAG stop codon"
FT	Misc-difference 208	/note= "Encoded by TAA stop codon"
FT	Misc-difference 234	/note= "Encoded by TAA stop codon"
FT	Misc-difference 245	/note= "Encoded by TGA stop codon"
FT	Misc-difference 253	/note= "Encoded by TGA stop codon"
FT	Misc-difference 269	/label= Encoded by TAA stop codon
FT	Misc-difference 295	/note= "Encoded by TAA stop codon"
FT	Misc-difference 298	/note= "Encoded by TAG stop codon"
FT	Misc-difference 323	/note= "Encoded by TGA stop codon"
FT	Misc-difference 324	/note= "Encoded by TGA stop codon"
FT	Misc-difference 331	/note= "Encoded by TGA stop codon"
FT	Misc-difference 341	/note= "Encoded by TAA stop codon"
FT	Misc-difference 344	/note= "Encoded by TAA stop codon"
FT	Misc-difference 356	/note= "Encoded by TAA stop codon"
FT	Misc-difference 358	/note= "Encoded by TGA stop codon"
FT	Misc-difference 413	/note= "Encoded by TAA stop codon"
FT	Misc-difference 469	/note= "Encoded by TGA stop codon"
FT	Misc-difference 486	/note= "Encoded by TAA stop codon"
FT	Misc-difference 501	/note= "Encoded by TGA stop codon"
FT	Misc-difference 523	/note= "Encoded by TGA stop codon"
FT	Misc-difference 532	/note= "Encoded by TGA stop codon"
FT	Misc-difference 586	/note= "Encoded by TGA stop codon"
FT	Misc-difference 593	/note= "Encoded by TGA stop codon"
FT	Misc-difference 596	/note= "Encoded by TGA stop codon"
FT	Misc-difference 632	/note= "Encoded by TAA stop codon"

FT	Misc-difference 640	/note= "Encoded by TGA stop codon"
FT	Misc-difference 681	/note= "Encoded by TGA stop codon"
FT	Misc-difference 695	/note= "Encoded by TAA stop codon"
FT	Misc-difference 710	/note= "Encoded by TGA stop codon"
FT	Misc-difference 717	/note= "Encoded by TGA stop codon"
FT	Misc-difference 731	/note= "Encoded by TGA stop codon"
FT	Misc-difference 747	/note= "Encoded by TGA stop codon"
FT	Misc-difference 756	/note= "Encoded by TGA stop codon"
FT	Misc-difference 764	/note= "Encoded by TAA stop codon"
FT	Misc-difference 775	/note= "Encoded by TAG stop codon"
FT	Misc-difference 780	/note= "Encoded by TGA stop codon"
FT	Misc-difference 804	/note= "Encoded by TGA stop codon"
FT	Misc-difference 807	/note= "Encoded by TAA stop codon"
FT	Misc-difference 813	/note= "Encoded by TAA stop codon"
FT	Misc-difference 854	/note= "Encoded by TAA stop codon"
FT	Misc-difference 860	/note= "Encoded by TAA stop codon"
FT	Misc-difference 866	/note= "Encoded by TAA stop codon"
FT	Misc-difference 875	/note= "Encoded by TAA stop codon"
FT	Misc-difference 877	/note= "Encoded by TGA stop codon"
FT	Misc-difference 882	/note= "Encoded by TGA stop codon"
FT	Misc-difference 887	/note= "Encoded by TGA stop codon"
FT	Misc-difference 891	/note= "Encoded by TGA stop codon"
FT	Misc-difference 910	/note= "Encoded by TAA stop codon"
FT	Misc-difference 916	/note= "Encoded by TGA stop codon"
FT	Misc-difference 942	/note= "Encoded by TGA stop codon"
FT	Misc-difference 975	/note= "Encoded by TAA stop codon"
FT	Misc-difference 976	/note= "Encoded by TGA stop codon"
FT	Misc-difference 981	/note= "Encoded by TGA stop codon"
FT	Misc-difference 997	/note= "Encoded by TGA stop codon"
FT	Misc-difference 1013	/note= "Encoded by TAA stop codon"
FT	Misc-difference 1020	/note= "Encoded by TGA stop codon"
FT	Misc-difference 1026	/note= "Encoded by TGA stop codon"
FT	Misc-difference 1028	/note= "Encoded by TGA stop codon"
FT	Misc-difference 1044	/note= "Encoded by TAA stop codon"
FT	Misc-difference 1070	/note= "Encoded by TGA stop codon"
FT	Misc-difference 1099	/note= "Encoded by TGA stop codon"

```
FT Misc-difference 1102 /note- "Encoded by TGA stop codon"
FT Misc-difference 1114 /note- "Encoded by TGA stop codon"
FT Misc-difference 1118 /note- "Encoded by TGA stop codon"
FT Misc-difference 1127 /note- "Encoded by TGA stop codon"
FT Misc-difference 1130 /note- "Encoded by TGA stop codon"
FT Misc-difference 1143 /note- "Encoded by TGA stop codon"
FT Misc-difference 1159 /note- "Encoded by TGA stop codon"
FT Misc-difference 1161 /note- "Encoded by TGA stop codon"
FT Misc-difference 1167 /note- "Encoded by TGA stop codon"
FT Misc-difference 1172 /note- "Encoded by TGA stop codon"
FT Misc-difference 1179 /note- "Encoded by TGA stop codon"
FT Misc-difference 1212 /note- "Encoded by TGA stop codon"
FT Misc-difference 1230 /note- "Encoded by TGA stop codon"
FT Misc-difference 1255 /note- "Encoded by TGA stop codon"
FT Misc-difference 1260 /note- "Encoded by TGA stop codon"
FT Misc-difference 1267 /note- "Encoded by TGA stop codon"
FT Misc-difference 1273 /note- "Encoded by TGA stop codon"
FT Misc-difference 1305 /note- "Encoded by TGA stop codon"
FT Misc-difference 1313 /note- "Encoded by TGA stop codon"
FT Misc-difference 1334 /note- "Encoded by TGA stop codon"
FT Misc-difference 1344 /note- "Encoded by TGA stop codon"
FT Misc-difference 1517 /note- "Encoded by TGA stop codon"
FT Misc-difference 1529 /note- "Encoded by TGA stop codon"

Query Match 19.5%; Score 74.5; DB 21; Length 1856;
Best Local Similarity 22.8%; Pred. No. 1.1;
Matches 21; Conservative 10; Mismatches 34; Indels 27; Gaps 3;

QY 2 DDDDDR-----RGHSIQOCYQRCRQRPRTSHARC-VQECRDDQOQHGRHEQ 49
Db 477 dqhderswhxkggrhpgpgrpgeaexklsalyedprsvrhaksgcpxrtdsgyghxqgmd 536
QY 50 EEOG-----RGRGMHGEEREE 66
Db 537 hvrgcktgcsaigentprkhtsrwshesrtqse 568

RESULT 10
W62838 standard; Protein; 605 AA.
XX W62838;
XX
XX 27-OCT-1998 (first entry)
XX
XX Glycine max antimicrobial protein.
XX
XX antimicrobial protein; infestation; control.
XX
XX Glycine max.
XX
XX WO9827805-A1.
XX
```

```
PD 02-JUL-1998.
XX
XX 22-DEC-1997; 97MO-AU00874.
XX
XX 20-DEC-1996; 96AU-0004275.
XX
XX (REFR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
XX
XX Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;
XX WPI; 1998-377279/32.
XX
XX Novel anti-microbial protein from e.g. Macadamia integrifolia -
XX useful for controlling microbial infestations of plants or mammals
XX Claim 1; Page 63-65; 96pp; English.
XX
XX "The sequence is that of an antimicrobial protein which can
XX be used to control microbial infestations in plants and mammalian
XX animals.
XX
XX Sequence 605 AA;

Query Match 19.1%; Score 73; DB 19; Length 605;
Best Local Similarity 30.3%; Pred. No. 0.47;
Matches 20; Conservative 6; Mismatches 14; Indels 26; Gaps 3;

QY 15 QCVORCQGERPRY-----SHARC-----VQEC-----RDDQOQHGRHEQ 48
Db 37 kclqscnsrdsyngqacharcmlkvekecegeglprprpqrphetepqpgxeed 96
QY 49 EEOGR 54
Db 97 edeqpr 102

RESULT 11
Y40999
ID Y40999 standard; Protein; 605 AA.
AC Y40999;
XX
XX 06-DEC-1999 (first entry)
XX
XX Soybean beta-conglycinin protein sequence.
XX
XX Peanut; allergen; Ara H 1; IgE; immunoglobulin E; epitope; Ara h 3;
XX allergic reaction; soybean; beta-conglycinin.
XX
XX Glycine max.
XX
XX WO9945961-A1.
XX
XX 16-SEP-1999.
XX
XX 12-MAR-1999; 99WO-US05494.
XX
XX 12-MAR-1998; 98US-0077763.
XX 11-MAR-1999; 99US-0077763.
XX
XX (UYAR-) UNIV ARKANSAS.
XX
XX Burke W, Helm RM, Cockrell G, Bannon GA, Stanley JS, Shin DS;
XX Sampson H, Compadre CM, Huang SK, Maleki SJ, Kopper RA;
XX WPI; 1999-551218/46.
XX
XX Tertiary structure of peanut allergen Ara h 1 for protection of a host
XX animal from allergic reaction -
XX
XX Disclosure; Fig 33A-B; 193pp; English.
XX
```

The invention provides a tertiary structure for the peanut allergen Ara H 1. The Ara H 1 allergen is found to contain 23 linear IgE-binding epitopes. The invention also provides an isolated recombinant peanut allergen designated Ara h 3 and a nucleotide molecule encoding the peanut allergen Ara h 3. Molecules of the invention are used to protect a host animal from allergic reaction, particularly using a modified allergen which is less reactive with IgE. The invention may also be used to ensure that the allergen is not introduced into genetically modified food. The present sequence represents a soybean beta-conglycinin protein sequence.

Sequence 605 AA:

Query Match	19.1%	Score 73:	DB 20:	Length 605:
Best Local Similarity	30.3%	Pred. No.	0.47:	
Matches 20:	Conservative 6:	Mismatches 14:	Indels 26:	Gaps 3:

```

QY 15 QCVDRCRQERRRY-----SHARC-----VQEC-----RDDQDQHGRRHQ 48
      ::: | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 37 kclqscnsrdsyngacharcnlkvekeockgypiprprpqhperpqgqkeed 96

```

QY	49	EEQGR	54
		1:111	
Db	97	edeqpr	102

RESULT	12
W62831	
ID	W62831 standard; Protein; 525 AA
XX	
AC	W62831;
XX	
DT	27-OCT-1998 (first entry)

DE	Theobroma cacao antimicrobial protein.
XX	
KM	antimicrobial protein; infestation; control
XX	
OS	Theobroma cacao.

XX		
PN	WO9827805-A1.	
XX		
XX	02-JUL-1998.	
PD		
XX		
PF	22-DEC-1997;	97WO-AU00874.
XX		
PR	20-DEC-1996;	96AU-0004275

PA (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.

PI Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP,
XX
DR WPI; 1998-377279/32.
22

PT Novel anti-microbial protein from e.g. *Mecadamia integrifolia* -
 PT useful for controlling microbial infestations of plants or mammals
 XX
 PS Claim 1: Page 47-49; 96pp; English.

CC The sequence is that of an antimicrobial protein which can
CC be used to control microbial infestations in plants and mammalian
CC animals.

525 AA;
Sequence

Query Match	18.8%	Score 72	DB 19	Length 525
Best Local Similarity	28.6%	Pred. No.	0.53	
Matches 22	Conservative	8	Mismatches 23	Indels 24
				Gaps 4

```

QY      14 00VQ0RCR0E0RPRYSH-----ARCVOEC--RDDQ0QHGRHQ-----EEEOG 53
      : 11 111 : | 1:1 | : 11 : 11 | 1:1:1

```

Db 60 egcaqrcreykeqqrqgeeelq,qyqqcgr,rceqeqqqr,eqqqrckweykeqerg 119
QY 54 KGRGWH----GEGERE 66
 :| | |
Db 120 ehennyhnhkknrrseeee 136

RESULT 13
R20181
ID R20181 standard; Protein; 566 AA.

DT 16-APR-1992 (first entry)
 XX

DE Sequence encoded by 67 kD T. cacao protein cDNA.
xy

KW Cocoa; flavour; seed storage protein.
YX

OS Theobroma cacao.
XY

PN WO9119801-A.
XY

PD 26-DEC-1991.
XY

PF 07-JUN-1991; 91WO-GB00914.
YY

PR 11-JUN-1990; 90GB-0013016.
XY

PA (MRSC) MARS UK LTD.
XY

PI Spencer ME, Hodge R, Deakin EA, Ashton S;
YX

DR WPI; 1992-024418/03.
DR N-PSDB; Q20377.

PT Recombinant cocoa proteins - are responsible for flavour in cocoa
PT beans and produced in large quantities using yeast and bacterial
PT expression vectors

PS Claim 4; Fig 2; 59pp; English.

The inventors claim a 67 kD and 31 kD T. cacao protein, and fragments, and encoding DNAs. The 47 kD and 31 kD proteins are detected from the 67 kD precursor. T. cacao protein cDNA was detected in a cDNA library prepared from immature cocoa beans RNA using a probe based on the AA sequence of a CNBR peptide common to the 47 kD and 31 kD polypeptides. Homology searches revealed close homologs between the 67 kD polypeptide and the vicilins, which are seed storage proteins.

50 Sequence 566 AA;

Query Match	18.8%	Score 72:	DB 13;	Length 566;
Best Local Similarity	28.6%;	Pred. No.	0.58;	
Matches 22;	Conservative	8;	Mismatches 23;	Indels 24;
				Gaps 4;

```
QY      14 QQCVCORCRDERPRYSH-----ARCVCFC-RDDQQQHGRHQ-----EEEG 53  
        : :: ||| : | :||| :||| :||| :|||  
Db      60 egcceqrccrcrkykeqgrqqeeelqrryqcqgrrcgeggqgrrcqqcqrlkewegykcgerr 119
```

QY 54 RGRGWH---GEGERHE 66

Db 120 ehenyhnhkkrseeee 1.36

RESULT 14
Y76583
ID Y76583 standard; Protein; 156 AA.

AC Y76583;
XX

```

DT 10-APR-2000 (first entry)
XX
XX Human ovarian tumor EST fragment encoded protein 79.
DE
XX Expressed sequence tag; EST; human; ovarian tumor; anticancer;
KW gene therapy; treatment.
XX
OS Homo sapiens.
XX
XX DE19817557-A1.
XX
XX 21-OCT-1999.
XX
XX 09-APR-1998; 98DE-1017557.
XX
XX 09-APR-1998; 98DE-1017557.
XX
XX 09-APR-1998; 98DE-1017557.
XX
XX (META-) METAGEN GES GENOMFORSCHUNG MBH.
XX
XX Rosenthal A, Specht T, Hinzmann B, Schmitt A, Pilarsky C, Dahl E;
XX
XX WPI; 1999-591920/51.
XX
XX N-PSDB; 277484.
XX
XX New nucleic acid sequences expressed in ovarian, and some other, cancer
XX
XX PT tissues, and derived polypeptides, for treatment of ovarian cancer and
XX
XX PT identification of therapeutic agents
XX
XX PS Claim 25; Page 276; 310pp; German.
XX
XX This invention describes novel nucleic acid (cDNA) sequences (A) which
XX
XX CC have anticancer activity and are highly expressed in ovarian tumor
XX
XX CC tissue (and some also in testis and breast cancer tissue). The products
XX
XX CC of the invention can be used for gene therapy. (A) are used (1) for
XX
XX CC recombinant expression of polypeptides (B) and (1i) to isolate complete
XX
XX CC genes. (B) are used (1) to identify agents suitable for treatment of
XX
XX CC ovarian cancer: (1i) directly for treating this form of cancer (including
XX
XX CC expression from gene therapy vectors) and (1ii) for generation of
XX
XX CC specific antibodies. (A) are identified by assembling ESTs (expressed
XX
XX CC sequence tags) from a particular tissue type before comparison of
XX
XX CC expression patterns. This allows a significantly longer fragment of the
XX
XX CC gene to be revealed, so should reduce the number of failures associated
XX
XX CC with the fact that ESTs from different libraries may represent different
XX
XX CC parts of the same unknown gene, distorting the estimated frequency of
XX
XX CC occurrence in a particular tissue. Y76505-Y7638 represent protein
XX
XX CC fragments encoded by the human ovarian tumor cDNA library derived EST
XX
XX CC fragments represented in 277450-277572.
XX
XX SO Sequence 156 AA;

Query Match 18.3%; Score 70; DB 20; Length 156;
Best Local Similarity 28.6%; Pred. No. 0.24;
Matches 18; Conservative 11; Mismatches 28; Indels 6; Gaps 2;

OY 7 RRGSHLQCCVQRCRQRPYSHA----RCVQECRDDQOQHGHDEEGRGKWHGEG 62
   | | | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 70 rsgskartpqlflgqlqfghgncvrcwlaqarehpgqgaqseegq--egq 127
   | | | | | | | | | | | | | | | | | | | | | | | | | | | |

OY 63 ERE 65
   : |
DB 128 qee 130

RESULT 15
Y32218
ID Y32218 standard; Protein: 1122 AA.
XX
XX Y32218;
XX
XX 15-FEB-2000 (first entry)
XX
XX DT Drosophila PER B protein.
DE

```

```

XX
XX PER B; period; perl gene; transcription factor; circadian rhythm;
KW jet lag; sleep disorder; depression; seasonal affective disorder;
KW fertility; therapy.
XX
XX Drosophila melanogaster.
XX
XX PN W09957137-A1.
XX
XX PD 11-NOV-1999.
XX
XX PF 06-MAY-1999; 99MO-US10072.
XX
XX PR 07-MAY-1998; 98US-0084610.
XX
XX PA (HARD ) HARVARD COLLEGE.
XX
XX PI Weitz CJ, Gekakis N, Staknis D;
XX
XX DR WPI; 2000-052938/04.
XX
XX PT Novel heterodimeric composition for identifying modulators used in
XX
XX PT diagnosing and treating circadian clock disruption disorders
XX
XX PS Disclosure; Fig 31; 96pp; English.
XX
XX This sequence represents the Drosophila melanogaster PER protein
XX
XX CC PER B. Splice variants are PER A (see Y32217) and PER C (see
XX
XX CC Y32219). PER forms a heteromeric complex together with TIM (see
XX
XX CC Y32221) protein and has a biological activity which inhibits
XX
XX CC transcription of the perl gene when the CLOCK protein is present in
XX
XX CC combination with BMAL1 protein (see Y32209). The invention is
XX
XX CC based on the discovery of the transcriptional mechanism regulating
XX
XX CC genes responsible for the establishment and/or maintenance of the
XX
XX CC circadian clock, and provides an assay for novel drugs aimed at
XX
XX CC restoration of a normal circadian cycle, the drugs being modulators
XX
XX CC of BMAL1-CLOCK-mediated transcription of E-box-linked genes. The
XX
XX CC drugs are used to treat conditions such as jet lag, sleep disorders,
XX
XX CC depression (seasonal affective disorder) and infertility. The
XX
XX CC invention also provides BMAL1 and CLOCK proteins with which to
XX
XX CC stimulate the transcription of an E-box-linked gene which regulates
XX
XX CC the circadian clock.
XX
XX SO .Sequence 1122 AA;

Query Match 18.3%; Score 70; DB 21; Length 1122;
Best Local Similarity 36.2%; Pred. No. 2.2;
Matches 21; Conservative 9; Mismatches 14; Indels 14; Gaps 4;

OY 22 QERPRYSHARVOE--CRDDQ--OOHGRH-----EQEEOGKRGKWHGGEREE 66
   : | | | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 103 eekprpsgfcvqqlcrtelqdgqhgndhsepqlaq1qgeeedqs--gsesadrlve 159
   : | | | | | | | | | | | | | | | | | | | | | | | | | | | |

Search completed: March 1, 2001, 16:09:35
Job time: 1327 sec

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